

Refine Search

Search Results -

Terms	Documents
L3 and (lung adj tumor)	256

Database:

US Pre-Grant Publication Full-Text Database
 US Patents Full-Text Database
 US OCR Full-Text Database
 EPO Abstracts Database
 JPO Abstracts Database
 Derwent World Patents Index
 IBM Technical Disclosure Bulletins

Search:

L4

Refine Search

Recall Text

Clear

Interrupt

Search History

DATE: Friday, June 09, 2006 [Printable Copy](#) [Create Case](#)

<u>Set Name</u> side by side	<u>Query</u>	<u>Hit Count</u>	<u>Set Name</u> result set
<i>DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=OR</i>			
<u>L4</u>	L3 and (lung adj tumor)	256	<u>L4</u>
<u>L3</u>	PRO831	261	<u>L3</u>
<i>DB=USPT; PLUR=YES; OP=OR</i>			
<u>L2</u>	PRO831	22	<u>L2</u>
<u>L1</u>	6951737.pn.	1	<u>L1</u>

END OF SEARCH HISTORY

SCORE Search Results Details for Application 10063534 and Search Result us-10-063-534-30.rag.

[Score Home](#) [Retrieve Application](#) [SCORE System](#) [SCORE](#) [Comments /](#)
[Page](#) [List](#) [Overview](#) [FAQ](#) [Suggestions](#)

This page gives you Search Results detail for the Application 10063534 and Search Result us-10-063-534-30.rag.

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GenCore version 5.1.9
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OM protein - protein search, using sw model

Run on: June 6, 2006, 21:30:04 ; Search time 200 Seconds
(without alignments)
166.884 Million cell updates/sec

Title: US-10-063-534-30
Perfect score: 386
Sequence: 1 MLLLTLLLLLLLLKGSCLW.....PAMTCSQAQPRGEGEKVGDG 73

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 2589679 seqs, 457216429 residues

Total number of hits satisfying chosen parameters: 2589679

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : A_Geneseq_8:*
1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
5: geneseqp2002s:*
6: geneseqp2003as:*
7: geneseqp2003bs:*
8: geneseqp2004s:*
9: geneseqp2005s:*
10: geneseqp2006s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query		DB	ID	Description
		Match	Length			
1	386	100.0	73	3	AA999346	Aay99346 Human PRO
2	386	100.0	73	4	AAB66095	Aab66095 Protein o
3	386	100.0	73	4	AAB87540	Aab87540 Human PRO
4	386	100.0	73	5	ABG95865	Abg95865 Human sec
5	386	100.0	73	6	ABU90890	Abu90890 Novel hum
6	386	100.0	73	6	ABO33949	Abo33949 Human sec
7	386	100.0	73	6	ABU71966	Abu71966 Novel hum
8	386	100.0	73	6	ABU71520	Abu71520 Human sec
9	386	100.0	73	6	ABU72301	Abu72301 Human PRO
10	386	100.0	73	6	ABU90974	Abu90974 Human PRO
11	386	100.0	73	6	ABO27295	Abo27295 Human sec
12	386	100.0	73	6	ABU92490	Abu92490 Human sec
13	386	100.0	73	6	ABU81160	Abu81160 Human sec
14	386	100.0	73	6	ABO53275	Abo53275 Novel hum
15	386	100.0	73	6	ABU98277	Abu98277 Novel hum
16	386	100.0	73	6	ABU89282	Abu89282 Novel hum
17	386	100.0	73	6	ABU82489	Abu82489 Novel hum
18	386	100.0	73	6	ABU96453	Abu96453 Human PRO

19	386	100.0	73	6	ABU72123	Abu72123	Human	PRO
20	386	100.0	73	6	ADB17087	Adb17087	Human	tra
21	386	100.0	73	6	ABO44253	Abo44253	Human	sec
22	386	100.0	73	6	ABO33588	Abo33588	Novel	hum
23	386	100.0	73	6	ADA19892	Ada19892	Novel	hum
24	386	100.0	73	6	ADB17275	Adb17275	Human	tra
25	386	100.0	73	6	ADA20064	Ada20064	Novel	hum
26	386	100.0	73	6	ABO34181	Abo34181	Human	sec
27	386	100.0	73	6	ADA00361	Ada00361	Human	sec
28	386	100.0	73	7	ABO44441	Abo44441	Human	sec
29	386	100.0	73	7	ABO33465	Abo33465	Novel	hum
30	386	100.0	73	7	ADB85603	Adb85603	Novel	hum
31	386	100.0	73	7	ADB68282	Adb68282	Human	PRO
32	386	100.0	73	7	ADB68089	Adb68089	Human	PRO
33	386	100.0	73	7	ADB90906	Adb90906	Novel	hum
34	386	100.0	73	7	ADC06986	Adc06986	Human	PRO
35	386	100.0	73	7	ADC17891	Adc17891	Human	PRO
36	386	100.0	73	7	ADC17165	Adc17165	Mammalian	
37	386	100.0	73	7	ADC14863	Adc14863	Novel	hum
38	386	100.0	73	7	ADC52358	Adc52358	Novel	hum
39	386	100.0	73	7	ADD70537	Add70537	Human	sec
40	386	100.0	73	7	ADD39614	Add39614	Human	sec
41	386	100.0	73	7	ADD70060	Add70060	Human	sec
42	386	100.0	73	7	ADD36034	Add36034	Novel	hum
43	386	100.0	73	7	ADD38181	Add38181	Human	sec
44	386	100.0	73	7	ADD39137	Add39137	Human	sec
45	386	100.0	73	7	ADD38660	Add38660	Human	sec

ALIGNMENTS

RESULT 1

AAY99346

ID AAY99346 standard; protein; 73 AA.

XX

AC AAY99346;

XX

DT 08-AUG-2000 (first entry)

XX

DE Human PRO831 (UNQ471) amino acid sequence SEQ ID NO:22.

XX

KW Human; PRO polypeptide; membrane bound protein; receptor; diagnosis;
transmembrane; secretion; immunoadhesion; pharmaceutical; screening.

XX

OS Homo sapiens.

XX

PN WO200012708-A2.

XX

PD 09-MAR-2000.

XX

PF 01-SEP-1999; 99WO-US020111.

XX

PR 01-SEP-1998; 98US-0098716P.

PR 01-SEP-1998; 98US-0098749P.

PR 01-SEP-1998; 98US-0098750P.

PR 02-SEP-1998; 98US-0098803P.

PR 02-SEP-1998; 98US-0098821P.

PR 02-SEP-1998; 98US-0098843P.

PR 09-SEP-1998; 98US-0099536P.

PR 09-SEP-1998; 98US-0099596P.

PR 09-SEP-1998; 98US-0099598P.

PR 09-SEP-1998; 98US-0099602P.

PR 09-SEP-1998; 98US-0099642P.

PR 10-SEP-1998; 98US-0099741P.

PR 10-SEP-1998; 98US-0099754P.

PR 10-SEP-1998; 98US-0099763P.

PR 10-SEP-1998; 98US-0099792P.

PR 10-SEP-1998; 98US-0099808P.

PR 10-SEP-1998; 98US-0099812P.

PR 10-SEP-1998; 98US-0099815P.

PR 10-SEP-1998; 98US-0099816P.

PR 15-SEP-1998; 98US-0100385P.

PR 15-SEP-1998; 98US-0100388P.

PR 15-SEP-1998; 98US-0100390P.

PR 16-SEP-1998; 98US-0100584P.

PR 16-SEP-1998; 98US-0100627P.

PR 16-SEP-1998; 98US-0100661P.

PR 16-SEP-1998; 98US-0100662P.

PR 16-SEP-1998; 98US-0100664P.

PR 17-SEP-1998; 98US-0100683P.

PR 17-SEP-1998; 98US-0100684P.

PR 17-SEP-1998; 98US-0100710P.

PR 17-SEP-1998; 98US-0100711P.

PR 17-SEP-1998; 98US-0100919P.

PR 17-SEP-1998; 98US-0100930P.

PR 18-SEP-1998; 98US-0100848P.

PR 18-SEP-1998; 98US-0100849P.

PR 18-SEP-1998; 98US-0101014P.

PR 18-SEP-1998; 98US-0101068P.

PR 18-SEP-1998; 98US-0101071P.

SCORE Search Results Details for Application 10063534 and Search Result us-10-063-534-30.olig.rag.

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OM protein - protein search, using sw model

Run on: June 6, 2006, 21:06:03 ; Search time 195 Seconds
(without alignments)
171.163 Million cell updates/sec

Title: US-10-063-534-30
Perfect score: 73
Sequence: 1 MLLLTLLLLLLLLKGSCLW.....PAMTCSQAQPRGEGEKVGDG 73

Scoring table: OLIGO
Gapop 60.0 , Gapext 60.0

Searched: 2589679 seqs, 457216429 residues

Word size : 6

Total number of hits satisfying chosen parameters: 27796

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : A_Geneseq_8:*

- 1: geneseqp1980s:*
- 2: geneseqp1990s:*
- 3: geneseqp2000s:*
- 4: geneseqp2001s:*
- 5: geneseqp2002s:*
- 6: geneseqp2003as:*
- 7: geneseqp2003bs:*
- 8: geneseqp2004s:*
- 9: geneseqp2005s:*
- 10: geneseqp2006s:*

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42	73	100.0	73	7	ADD36034	Add36034 Novel hum
43	73	100.0	73	7	ADD38181	Add38181 Human sec
44	73	100.0	73	7	ADD39137	Add39137 Human sec
45	73	100.0	73	7	ADD38660	Add38660 Human sec

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PR 02-SEP-1998; 98US-0098843P.

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PR 09-SEP-1998; 98US-0099598P.

PR 09-SEP-1998; 98US-0099602P.

PR 09-SEP-1998; 98US-0099642P.

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PR 10-SEP-1998; 98US-0099763P.

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PR 10-SEP-1998; 98US-0099808P.

PR 10-SEP-1998; 98US-0099812P.

PR 10-SEP-1998; 98US-0099815P.

PR 10-SEP-1998; 98US-0099816P.

PR 15-SEP-1998; 98US-0100385P.

PR 15-SEP-1998; 98US-0100388P.

PR 15-SEP-1998; 98US-0100390P.

PR 16-SEP-1998; 98US-0100584P.

PR 16-SEP-1998; 98US-0100627P.

PR 16-SEP-1998; 98US-0100661P.

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PR 18-SEP-1998; 98US-0100848P.

PR 18-SEP-1998; 98US-0100849P.

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PR 18-SEP-1998; 98US-0101068P.

PR 18-SEP-1998; 98US-0101071P.